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## In the Claims:

- Please cancel claims 23 and 24 without prejudice.
- Please amend claims 15 22, 25, and 27 30 as follows:
  - (twice amended) The nucleotide sequence SEQ ID NO: 1 [from the Sequence Listing].
  - (twice amended) The nucleotide sequence SEQ ID NO: 2 [from the Sequence Listing].
  - 17. (twice amended) The nucleotide sequence SEQ ID NO: 3 [from the Sequence Listing].
  - 18. (twice amended) The nucleotide sequence SEQ ID NO: 4 [from the Sequence Listing].
  - 19. (twice amended) The nucleotide sequence SEQ ID NO: 5 [from the Sequence Listing].
  - 20. (twice amended) The nucleotide sequence SEQ ID NO: 6 [from the Sequence Listing].
  - 21. (twice amended) The nucleotide sequence SEQ ID NO: 7 [from the Sequence Listing].

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22. (twice amended) The nucleotide sequence SEQ ID NO: 8 [from the Sequence Listing].

25. A kit for the analysis of fungal infections with azole derivative-resistant

fungal strains, containing at least one nucleotide sequence[s] selected from

the group consisting of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3, SEQ

ID NO: 4, SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7[,] and SEQ ID NO:

8.

27. (once amended) A method for detecting azole derivative-resistant fungal cells

in clinical material, comprising the steps of:

a) extraction of fungus-specific nucleic acids from clinical material; and

b) hybridization of the fungus-specific nucleic acids with hybridization

probes which are directed against nucleic acid segments of azole derivative-

resistant fungal cells,

wherein detection of the hybridized probes detects azole derivative-resistant

cells,

wherein the hybridization probes are directed against a DNA segment from

the  $14-\alpha$ -lanosterol demethylase gene,

wherein between steps a) and b) a PCR reaction is performed in which

segments of the  $14-\alpha$ -lanosterol demethylase gene are amplified, and

wherein a primer for the PCR reaction is selected from the group consisting

of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3 and SEQ ID NO: 4.

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28. (once amended) A method for detecting azole derivative-resistant fungal cells

in clinical material., comprising the steps of:

a) extraction of fungus-specific nucleic acids from clinical material; and

b) hybridization of the fungus-specific nucleic acids with hybridization

probes which are directed against nucleic acid segments of azole derivative-

resistant fungal cells,

wherein detection of the hybridized probes detects azole derivative-resistant

cells,

wherein the hybridization probes are directed against a DNA segment from

the 14-α-lanosterol demethylase gene (ERG16 gene) of the species Candida

albicans,

wherein between steps a) and b) a PCR reaction is performed in which

segments of the 14-α-lanosterol demethylase gene are amplified, and

wherein a primer for the PCR reaction is selected from the group consisting

of SEQ ID NO: 1, SEQ ID NO: 2, SEQ ID NO: 3 and SEQ ID NO: 4.

29. (once amended) A method for detecting azole derivative-resistant fungal cells

in clinical material, comprising the steps of:

a) extraction of fungus-specific nucleic acids from clinical material; and

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b) hybridization of the fungus-specific nucleic acids with hybridization

probes which are directed against nucleic acid segments of azole derivative-

resistant fungal cells,

wherein detection of the hybridized probes detects azole derivative-resistant

cells,

wherein a [the] hybridization probe [probes] for step b) is selected from the

group consisting of SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7 and SEQ

ID NO: 8.

30. (once amended) A method for detecting azole derivative-resistant fungal cells

in clinical material, comprising the steps of:

a) extraction of fungus-specific nucleic acids from clinical material; and

b) hybridization of the fungus-specific nucleic acids with hybridization

probes which are directed against nucleic acid segments of azole

derivative-resistant fungal cells,

wherein detection of the hybridized probes detects azole derivative-resistant

cells,

wherein the hybridization probes are directed against a DNA segment from

the  $14-\alpha$ -lanosterol demethylase gene,

wherein between steps a) and b) a PCR reaction is performed in which

segments of the  $14-\alpha$ -lanosterol demethylase gene are amplified, and

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wherein a hybridization probe for step b) is selected from the group consisting of SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7 and SEQ ID NO: 8.

- Please add new claims 31 and 32, as follows:
  - 31. (new) The method of claim 27, wherein the primers are SEQ ID NO: 1 and 2, and the probes are SEQ ID NO: 5 and/or 6.
- 32. (new) The method of claim 27, wherein the primers are SEQ ID NO: 3 and 4, and the probes are SEQ ID NO: 7 and/or 8.
- Please note claims 10, 12 and 13 remain pending without amendment, as set forth below:
  - 10. The method of claim 27, wherein a hybridization probe for step b) is selected from the group consisting of SEQ ID NO: 5, SEQ ID NO: 6, SEQ ID NO: 7 and SEQ ID NO:8.
  - 12. The method of claim 29, wherein after hybridization, at least one washing step is performed at a temperature which is approximately 1°C less than the melting temperature (Tm) of the particular hybridization probe used.
- 13. The method of claim 30, wherein after hybridization, at least one washing step is performed at a temperature which is approximately l°C less than the melting temperature (Tm) of the particular hybridization probe used.